

SEQUENCE LISTING

<110> Lorenz, M., et al.

<120> A NOVEL P-SELECTIN GLYCOPROTEIN LIGAND (PSGL-1)
BINDING PROTEIN AND USES THEREFOR

<130> GFN-5380

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<150> 60/192,104

<151> 2000-03-24

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<170> PatentIn Ver. 2.0

<210> 1

<211> 951

<212> DNA

<213> Homo sapiens

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<222> (1)..(948)

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acc cag tgc acg gca agg acc cag cag gaa gca cca gcc act ggc ccc	96
Thr Gln Cys Thr Ala Arg Thr Gln Gln Glu Ala Pro Ala Thr Gly Pro	
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gac ctc ccg cac cca gga cct gac ggg cac tta gac aca cac agt ggc	144
Asp Leu Pro His Pro Gly Pro Asp Gly His Leu Asp Thr His Ser Gly	
35 40 45	
ctg agc tcc aac tcc agc atg acc acg cgg gag ctt cag cag tac tgg	192
Leu Ser Ser Asn Ser Ser Met Thr Thr Arg Glu Leu Gln Gln Tyr Trp	
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cag aac cag aaa tgc cgc tgg aag cac gtc aaa ctg ctc ttt gag atc	240
Gln Asn Gln Lys Cys Arg Trp Lys His Val Lys Leu Leu Phe Glu Ile	
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gct tca gct cgc atc gag gag aga aaa gtc tct aag ttt gtg gtg tac	288
Ala Ser Ala Arg Ile Glu Glu Arg Lys Val Ser Lys Phe Val Val Tyr	
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caa atc atc gtc atc cag act ggg agc ttt gac aac aac aag gcc gtc	336
Gln Ile Ile Val Ile Gln Thr Gly Ser Phe Asp Asn Asn Lys Ala Val	
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ctg gaa cgg cgc tat tcc gac ttc gcg aag ctc cag aaa gcg ctg ctg	384
Leu Glu Arg Arg Tyr Ser Asp Phe Ala Lys Leu Gln Lys Ala Leu Leu	
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Lys Thr Phe Arg Glu Glu Ile Glu Asp Val Glu Phe Pro Arg Lys His
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ctg act ggg aac ttc gct gag gag atg atc tgt gag cgt cgg cgc gcc 480
Leu Thr Gly Asn Phe Ala Glu Glu Met Ile Cys Glu Arg Arg Arg Ala
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ctg cag gag tac ctg ggc ctg ctc tac gcc atc cgc tgc gtg cgc cgc 528
Leu Gln Glu Tyr Leu Gly Leu Leu Tyr Ala Ile Arg Cys Val Arg Arg
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tcc cgg gag ttc ctg gac ttc ctc acg cgg ccg gag ctg cgc gag gct 576
Ser Arg Glu Phe Leu Asp Phe Leu Thr Arg Pro Glu Leu Arg Glu Ala
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ttc ggc tgc ctg cgg gcc ggc cag tac ccg cgc gcc ctg gag ctg ctg 624
Phe Gly Cys Leu Arg Ala Gly Gln Tyr Pro Arg Ala Leu Glu Leu Leu
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Leu Arg Val Leu Pro Leu Gln Glu Lys Leu Thr Ala His Cys Pro Ala
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gcc gcc gtc ccg gcc ctg tgc gcc gtg ctg ctg tgc cac cgc gac ctc 720
Ala Ala Val Pro Ala Leu Cys Ala Val Leu Leu Cys His Arg Asp Leu
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gac cgc ccc gcc gag gcc ttc gcg gcc gga gag agg gcc ctg cag cgc 768
Asp Arg Pro Ala Glu Ala Phe Ala Ala Gly Glu Arg Ala Leu Gln Arg
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Leu Gln Ala Arg Glu Gly His Arg Tyr Tyr Ala Pro Leu Leu Asp Ala
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Glu Arg Leu Glu Glu Ser Gln Leu Arg Arg Pro Thr Pro Arg Gly Ile
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 Gln Asn Gln Lys Cys Arg Trp Lys His Val Lys Leu Leu Phe Glu Ile
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 Ser Arg Glu Phe Leu Asp Phe Leu Thr Arg Pro Glu Leu Arg Glu Ala
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 Ala Ala Val Pro Ala Leu Cys Ala Val Leu Leu Cys His Arg Asp Leu
 225 230 235 240
 Asp Arg Pro Ala Glu Ala Phe Ala Ala Gly Glu Arg Ala Leu Gln Arg
 245 250 255
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 Met Val Arg Leu Ala Tyr Ala Leu Gly Lys Asp Phe Val Thr Leu Gln
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<223> Description of Artificial Sequence: primer

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